• Sample code: NUT18247AA

• Sample date: 26-06-2023



## Fagron NutriGen™

Patient report



• Sample code: NUT18247AA



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- 2. Genetic results overview
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### **Patient identification data**

1



Patient's name and last name — DEMO 2 WOMAN Structure

Date of Birth — 26-09-1953

Gender — woman

Sample type — Buccal mucosa

Sample code — NUT18247AA

Sample date — 04-03-2024

Report date — 26-06-2023

Doctor's name and last name — Nutrigen 3 Validation

Email address — nutrigen\_3\_validation@demo.com

Telephone — •

### LEGAL DISCLAMER

Fagron Genomics, S.L.U carries out genetic tests upon request by healthcare professionals, in relation to biological samples from patients obtained by the healthcare professional. Our tests do not replace a medical consultation, nor do they make up a diagnostic or treatment, nor should they be interpreted this way. Only healthcare professionals can interpret the results of said tests, based on their knowledge of the clinical records of the patients and other relevant factors and, under their responsibility, give a diagnostic or prescribe treatment to the patient. We decline all responsibility derived from the use and interpretation of the results of our tests by the solicitant healthcare professional. Fagron Genomics, S.L.U expressly reserves any legal actions in case of an inappropriate, negligent or incorrect use or interpretation of the results of our tests. It is the responsibility of the healthcare professional who requests a test to guarantee to the patient the appropriate genetic advice as foreseen by Law 14/2007, of 3rd July, of biomedical research. As Fagron Genomics, S.L.U does not have access to the personal identifiable information about the patient from whom the sample comes, it is the responsibility of the requesting healthcare professional to comply with the applicable data protection Laws and regulations.



Fagron Genomics, SRN: ES-MF-00001092 C/ de les Cosideres, 150 08226 Terrassa, Barcelona (Spain)







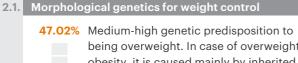


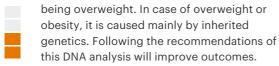
Product version: NutriGen<sup>TM</sup> 3.0 Report version: Rev.5 / 04-03-2024 08:21:44



• Sample date: **26-06-2023** 

### **Genetic results overview**





Genetic risk of overweight/obesity
Risk of rebound weight gain
Risk of increased BMI

Basal metabolic rate (burn calories at rest)
Weight loss capability during diet
interventions

#### 2.2. Behavioural genetics in food intake

73.79% Medium-low dysregulation of food intake behaviour. Slight predisposition to being overweight. In case of excessive quantity or compulsive intake, strategies to reduce anxiety should be considered.



### 2.3. Flavour sensitivities

99.67% Normal or average flavour sensitivity.



Bitter taste sensitivity
Salt sensitivity
Sweet flavour preference

#### 2.4. Fat metabolism

67.61% Moderate fat burning capacity. A healthy fat intake should not be a reason for being overweight.



### 2.5. Lipid metabolism

50.12% Moderately affected lipid metabolism.

Cholesterol and triglyceride levels should be reasonably normal on a balanced diet.



Predisposition to increased oxidation of

Risk of increased cholesterol LDL levels
Risk of unbalanced Triglycerides/HDL ratio

### 2.6. Carbohydrate metabolism

**24.92%** Highly negative carbohydrate metabolism:

Carbohydrate intake will lead to dysregulation in cholesterol levels and also to increased calorie and fat intake.

Eliminating refined carbohydrates is urgent; move to wholegrain carbohydrates and

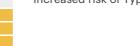
reduce the quantity.



Capability to digest starchy food Refined carbohydrate sensitivity Carbohydrates and HDL levels predisposition Carbohydrates and LDL levels

### 2.7. Glucose metabolism

**56.32%** Slightly affected glucose metabolism. Increased risk of Type-II diabetes.



Risk of increased glucose levels in plasma after fasting Risk of insulin resistance Risk of Type-II diabetes

#### 2.8. Efficacy of exercise

63.17% Medium efficacy of exercise to reduce body fat and regulate cholesterol levels. While dietary interventions may be needed, please do not forget to increase exercise. The benefits will be noticeable.



### **INDICATIONS**

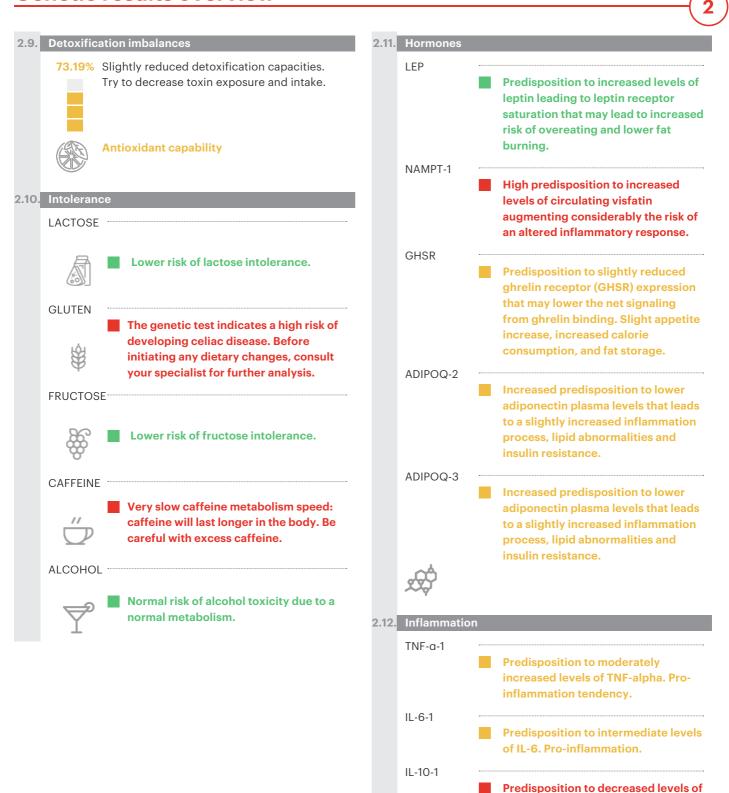
Positive effect Medium-positive effect Medium-negative effect Negative effect

\* These recommendations are based only in the analysis of your genetic test. Always seek the advice of your physician or other qualified health specialist before proceeding with any nutritional or dietary modifications.

• Sample date: **26-06-2023** 

• Date of the results: 30-06-2023

### **Genetic results overview**



Positive effect

Medium-positive effect

Medium-negative effect

Negative effect

the anti-inflammatory cytokine IL-

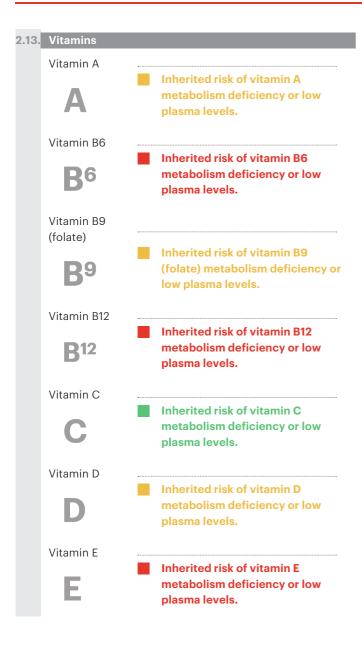
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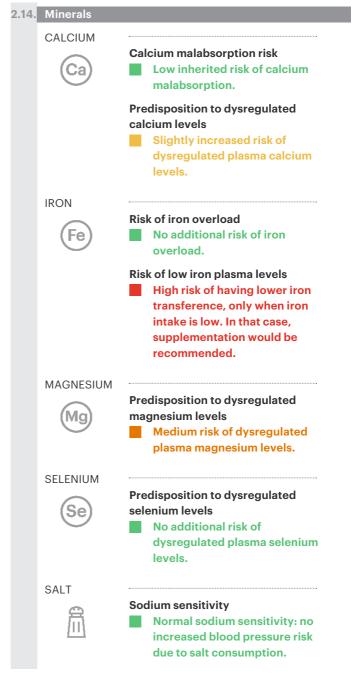


INDICATIONS

• Sample date: 26-06-2023

### **Genetic results overview**





#### INDICATIONS









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• Sample date: 26-06-2023

### **Genetic results overview**



### **Recommended nutritional plan**

3



The most effective diet for your patient, after the genetic analysis, would be

LOW IN CARBOHYDRATES
INTEGRATED NUTRITIONAL PLAN



### **Check the FOOD LIST**

recommended for you



### **Recommended supplements**



### The supplements recommended

to combat overweight and ageing are divided into 3 phases







### **INTESTINAL**

**SUPPLEMENTATION** 

Detoxific	catio	on
(oxidation)	30	days

### (conjugation) 30 days

### Transportation Excretion **30 days**

Prevention, maintaining optimal nutrition **3-6 months** 

- Vitamin E
- Vitamin B6 (Pyridoxine hydrochloride)
- Selenium (Selenium yeast)
- Brocophanus®
- Glutathione (Reduced glutathione)
- Resveratrol
- Silybin®
- Ubiqsome®
- Betacarotene
- Manganese
- Ubiquinol
- Zinc gluconate

- Magnesium
- Vitamin B9 (Methylfolate)
- Allyl ABG™(Allium sativum)
- Vitamin D3 (Cholecalciferol)
- CitrusiM®
- Brocophanus®
- Glutathione (Reduced glutathione)
- Resveratrol
- Alpha-Lipoic
   Acid (ALA)
- Silybin®
- Citrimax®
- Taurine

- Biointestil®
- · Lactobacillus salivarius
- Gutcare®
- · Lactobacillus lactis
- · Bifidobacterium longum
- Papain
- · Lactobacillus acidophilus
- Bifidobacterium infantis
- · Lactobacillus plantarum
- Bifidobacterium adolescentis
- Ginger dry extract
- · Bifidobacterium bifidum

- Magnesium
- Glucosamine sulfate
- Vitamin E
- SiliciuMax® powder
- Vitamin B6 (Pyridoxine hydrochloride)
- Vitamin B9 (Methylfolate)
- Vitamin B12
   (Cianocobalamin)
- Selenium (Selenium yeast)
- Vitamin D3
   (Cholecalciferol)
- CitrusiM®
- FitNox®
- Resveratrol

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### **Complete genetic results**



	2.1. Morphological genetics for	weight control			
	GENETIC RISK	MARKER	LOCUS	YOUR VARIANT	YOUR RESULT
)		MC4R-1	rs2229616	CC	
1	Genetic risk of overweight/obesity	SH2B1-2	rs7498665	AG	
	MEDIUM-LOW RISK OF	FTO-1	rs9939609	TT	
	OVERWEIGHT/OBESITY	FTO-2	rs1121980	AG	
		MC4R-2	rs17700633	GG	
	Risk of rebound weight gain	ADIPOQ	rs17300539	AG	
	MEDIUM-LOW REBOUND EFFECT	7,511 0 0	101700000	710	
	Risk of increased BMI	MC4R-3	rs12970134	GG	
	MEDIUM-LOW RISK OF	MC4R-4	rs17782313	CT	
	INCREASED BMI	SH2B1-1	rs4788102	AG	
	Basal metabolic rate (burn calories at rest)	FABP2	rs1799883	СТ	
	MEDIUM-LOW BURNER AT REST	LEPR-4	rs2025804	АА	
	Weight loss capability during				
	diet interventions	ACSL5	rs2419621	CC	
	SLOW WEIGHT LOSS				

### **INDICATIONS**





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• Sample date: 26-06-2023

### **Complete genetic results**

5

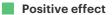
	2.2. Behavioural genetics in fo	ood intake			
	GENETIC RISK	MARKER	Locus	YOUR VARIANT	YOUR RESULT
)		COMT	rs4680	AG	
1	Appetite and anxiety risk	NMB	rs1051168	GT	
	INCREASED APPETITE AND	DRD2-1	rs1800497	AG	
	ANXIETY RISK	MC4R-1	rs2229616	CC	
		DRD2-2	rs6277	AG	
	Satiety: Feeling Full NORMAL SATIETY	FTO-1	rs9939609	TT	



2.3. Flavour sensitivities					
	GENETIC RISK	MARKER	LOCUS	YOUR VARIANT	YOUR RESULT
	Bitter taste sensitivity	TAS2R38-1	rs1726866	GG	
	NORMAL BITTER TASTE SENSITIVITY	TAS2R38-2	rs713598	CG	
	Salt sensitivity LOW SALT SENSITIVITY	ACE	rs4343	АА	
	Sweet flavour preference NORMAL SWEET PREFERENCE	SLC2A2	rs5400	GG	

### **INDICATIONS**





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• Sample date: 26-06-2023

### **Complete genetic results**



	2.4. Fat metabolism				
	GENETIC RISK	MARKER	LOCUS	YOUR VARIANT	YOUR RESULT
7,000	Response to monounsaturated fats (MUFAs) FAST MUFA METABOLISM	ADIPOQ	rs17300539	AG	
	Response to polyunsaturated fats (PUFAs)	PPAR-Y	rs1801282	CC	
	LOW PUFA METABOLISM	FADS1	rs174547	CC	
	Response to fat intake to improve the HDL levels  MEDIUM-HIGH BENEFITS TO IMPROVE HDL	LIPC	rs1800588	СТ	•

### **INDICATIONS**



Medium-positive effect

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### **Complete genetic results**



2.5. Lipid metabolism					
	GENETIC RISK	MARKER	LOCUS	YOUR VARIANT	YOUR RESULT
	Predisposition to reduced HDL levels	APOA5	rs662799	AA	
	REDUCED HDL LEVELS	СЕТР	rs5883	CC	
	Predisposition to increased levels of triglycerides HIGHLY INCREASED TRIGLYCERIDES	PPAR-Y	rs1801282	CC	•
	Predisposition to increased oxidation of LDL  NOT INCREASED LDL  OXIDATION	APOB-2	rs676210	АА	•
	Risk of increased cholesterol LDL	CELSR2	rs12740374	GT	
	levels	HNF1A	rs2650000	AC	
	INCREASED LDL LEVELS	LDLR	rs6511720	GG	
	MOREAGED EDE EEVEEG	ABCG8	rs6544713	СТ	
	Risk of unbalanced Triglycerides/HDL ratio SLIGHLTLY INCREASED TG/HDL RATIO	HMGCR	rs3846663	СТ	•

### **INDICATIONS**





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• Sample date: 26-06-2023

### **Complete genetic results**



	2.6. Carbohydrate metabolism					
	GENETIC RISK	MARKER	LOCUS	YOUR VARIANT	YOUR RESULT	
3	Capability to digest starchy food HIGHLY REDUCED STARCH DIGESTION	AMY1-AMY2 AMY1	rs11577390 rs4244372	CC AA		
	Refined carbohydrate sensitivity  NORMAL CARBOHYDRATE  SENSITIVITY	FABP2	rs1799883	СТ	•	
	Carbohydrates and HDL levels predisposition HIGH RISK OF HDL DYSREGULATION	KCTD10	rs10850219	GG		
	Carbohydrates and LDL levels HIGH RISK OF LDL DYSREGULATION	ММАВ	rs2241201	CG	•	

### **INDICATIONS**



Medium-positive effect

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### **Complete genetic results**

5

	2.7. Glucose metabolism				
	GENETIC RISK	MARKER	Locus	YOUR VARIANT	YOUR RESULT
	Risk of increased glucose levels in plasma after fasting	PLIN1	rs2289487	СТ	
	MEDIUM-LOW RISK OF HIGH GLUCOSE LEVELS	GHSR	rs490683	GC	
		PPAR-Y	rs1801282	CC	
	Risk of insulin resistance	ADIPOQ	rs17300539	AG	
	MEDIUM-LOW INSULIN RESISTANCE	TCF7L2-2	rs7903146	CC	
		FTO-1	rs9939609	TT	
		FTO-2	rs1121980	AG	
		PPAR-Y	rs1801282	CC	
		PLIN1	rs2289487	CT	
		TCF7L2-2	rs7903146	CC	
		FTO-1	rs9939609	TT	
	Risk of Type-II diabetes	MC4R-2	rs17700633	GG	
	MEDIUM-LOW DIABETES TYPE-II RISK	CDKN2A/B	rs10811661	СТ	
		KCNQ1	rs2237892	СТ	
		CDKN2A, CDKN2B	rs2383208	AG	
		CDKAL1	rs7756992	AG	
		TCF7L2-1	rs7901695	TT	

### **INDICATIONS**





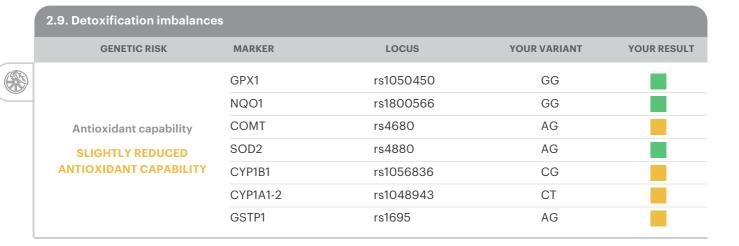
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### **Complete genetic results**



2.8. Efficacy of exercise				
GENETIC RISK	MARKER	LOCUS	YOUR VARIANT	YOUR RESULT
Benefits from endurance exercise for improving HDL levels	PPARD	rs2016520	CC	
HIGH BENEFITS FROM EXERCISE FOR IMPROVING HDL				
	FTO-1	rs9939609	TT	
Exercise to reduce body fat  MEDIUM-LOW BENEFIT FROM  EXERCISE TO REDUCE FAT	FTO-2	rs1121980	AG	
	LIPC	rs1800588	СТ	
	LEP	rs7799039	AA	



### **INDICATIONS**





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	2.10. Intolerance				
	GENETIC RISK	MARKER	LOCUS	YOUR VARIANT	YOUR RESULT
	Lactose intolerance risk	MCM6-1	rs182549	СТ	
	LOWER RISK OF LACTOSE INTOLERANCE	MCM6-2	rs4988235	AG	
(AB)		HLA-7	rs2187668	CT	
4		HLA-8	rs4639334	GG	
	Risk of celiac disease	HLA-2	rs2395182	TT	
	HIGHER RISK OF CELIAC DISEASE	HLA-4	rs4713586	AA	
	DIOLAGE	HLA-5	rs7454108	TT	
		HLA-6	rs7775228	TC	
<b>\$</b>	Fructose intolerance risk	ALDOB-1	rs1800546	CC	
	LOWER RISK OF FRUCTOSE INTOLERANCE	ALDOB-2	rs76917243	GG	
("	Caffeine metabolism	CYP1A1-1	rs2470893	CC	
	SLOW CAFFEINE METABOLIZER	CYP1A2	rs762551	CA	
7	Alcohol metabolism				
1	NORMAL ALCOHOL METABOLISM	ALDH2	rs671	GG	

### **INDICATIONS**

Negative effect

Medium-positive effect

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#### • Sample date: 26-06-2023

### **Complete genetic results**

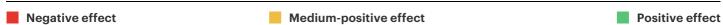
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	2.11. Hormones				
	GENETIC RISK	MARKER	LOCUS	YOUR VARIANT	YOUR RESULT
>	Leptin  LOW RISK OF DECREASED  LEPTIN LEVELS	LEP	rs7799039	АА	•
	Visfatin HIGH RISK OF INCREASED VISFATIN LEVELS	NAMPT-1	rs9770242	AA	
	Ghrelin LOW-INTERMEDIATE GHRELIN RECEPTOR (GHSR) EXPRESSION	GHSR	rs490683	GC	•
	Adiponectin  MEDIUM-LOW RISK OF  DECREASED ADIPONECTIN  LEVELS	ADIPOQ-2 ADIPOQ-3	rs1501299 rs2241766	GT GT	



	2.12. Inflammation				
	GENETIC RISK	MARKER	LOCUS	YOUR VARIANT	YOUR RESULT
5	TNF-a  MEDIUM-LOW RISK OF INCREASED TNF-a LEVELS	TNF-a-1	rs1800629	AG	•
	IL-6 MEDIUM-LOW RISK OF INCREASED IL-6 LEVELS	IL-6-1	rs1800795	CG	•
	IL-10 HIGH RISK OF DECREASED ANTIINFLAMATORY CYTOKINE IL-10 LEVELS	IL-10-1	rs1800896	TT	

### **INDICATIONS**



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### **Complete genetic results**

5

	2.13. Vitamins				
ĺ	GENETIC RISK	MARKER	Locus	YOUR VARIANT	YOUR RESULT
A	Vitamin A	BCMO1-1	rs12934922	TT	
	MEDIUM-LOW RISK OF VITAMIN A DEFICIENCY	BCMO1-2	rs7501331	CC	
B <sub>6</sub>	Vitamin B6				
	HIGH RISK OF VITAMIN B6 DEFICIENCY	NBPF3	rs4654748	CC	
B <sub>9</sub>	Vitamin B9 (folate)				
	MEDIUM-LOW RISK OF VITAMIN B9 (Folate) DEFICIENCY	MTHFR	rs1801133	GA	
B12	Vitamin B12				
	HIGH RISK OF VITAMIN B12 DEFICIENCY	FUT2	rs602662	GG	
C	Vitamin C	SLC23A2	rs1279683	GA	
	LOW RISK OF VITAMIN C DEFICIENCY	SLC23A1	rs33972313	CC	
D		GC	rs2282679	TT	
	Vitamin D	CYP2R1-2	rs10741657	AG	
	MEDIUM-LOW RISK OF VITAMIN	NADSYN1, DHCR7-1	rs12785878	GT	
	D DEFICIENCY	CYP2R1-3	rs2060793	AG	
		NADSYN1, DHCR7-2	rs3829251	GG	
E	Vitamin E	INTERGENIC	rs12272004	CC	
	HIGH RISK OF VITAMIN E DEFICIENCY	ZNF259, LOC100128347, APOA5, SIK3, BUD13	rs964184	CC	

### **INDICATIONS**

Negative effect

Medium-positive effect

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### **Complete genetic results**



	2.14. Minerals				
	GENETIC RISK	MARKER	LOCUS	YOUR VARIANT	YOUR RESULT
(Ca)	Calcium malabsorption risk	CYP2R1-1	rs10766197	GG	
	LOW RISK OF CALCIUM MALABSORPTION	GC	rs2282679	TT	
		DGKD	rs1550532	CG	
	Predisposition to dysregulated calcium levels  SLIGHTLY INCREASED RISK OF DYSREGULATED PLASMA CALCIUM LEVELS	CYP24A1	rs1570669	AA	
		CASR-1	rs17251221	AA	
		CASR-2	rs1801725	GG	
		CARS	rs7481584	AA	
		GCKR	rs780094	СТ	
Fe	Risk of iron overload  LOW RISK OF HEMOCHROMATOSIS	HFE	rs1800562	GG	•
	Risk of low iron plasma levels HIGH RISK OF DECREASED IRON LEVELS	TF-1	rs3811647	AA	
		TMPRSS6	rs4820268	AG	
		TF-2	rs8177253	TT	
Mg	Predisposition to dysregulated magnesium levels	CASR-1	rs17251221	AA	
		TRPM6	rs11144134	TT	
	MEDIUM-HIGH RISK OF DYSREGULATED MAGNESIUM LEVELS	SHROOM3	rs13146355	GG	
		DCDC5	rs3925584	CT	
		MUC1	rs4072037	СТ	
Se	Predisposition to dysregulated selenium levels	AGA	rs1395479	CC	
	NO ADDITIONAL RISK OF DYSREGULATED SELENIUM LEVELS	SLC39A11	rs891684	GG	-
	Sodium sensitivity LOW SODIUM SENSITIVITY	ACE	rs4343	AA	

### **INDICATIONS**

Negative effect

Medium-positive effect

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### **Complete genetic results**

5

	2.15. Effectiveness of diets						
	GENETIC RISK	MARKER	LOCUS	YOUR VARIANT	YOUR RESULT		
9	Efficacy of low calorie diets  VERY LOW EXPECTED BENEFIT  FROM LOW-CALORIE DIET	PPAR-Y	rs1801282	CC			
		ADIPOQ	rs17300539	AG			
		LEPR-1	rs1805134	TT			
		ACSL5	rs2419621	CC			
		ADRB2	rs1042714	CC			
	Efficacy of low carbohydrate diets						
	MEDIUM-HIGH EXPECTED BENEFIT FROM LOW- CARBOHYDRATE DIET	KCTD10	rs10850219	GG			
		ММАВ	rs2241201	CG	•		
		PPAR-Y	rs1801282	CC			
	Efficacy of low fat diets  MEDIUM-LOW EXPECTED BENEFIT FROM LOW-FAT DIET	GHSR	rs490683	GC			
		APOA2	rs5082	AG			
		SH2B1-2	rs7498665	AG			
		TCF7L2-2	rs7903146	CC			
		FTO-1	rs9939609	TT			

### **INDICATIONS**





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### Methodology



### How were the genetic variants selected and evaluated?

This test was developed by a multidisciplinary team of medical doctors, geneticists, and programmers, following highest quality standards. In particular, an expert team specialized in the curation of genetic variants reviewed each variant to ensure that selection, interpretation and impact of variants in the algorithms are based on the highest scientific evidence.

The following selection criteria were applied for classifying genetic variants:

**Level 1A:** Annotation for a variant in medical society-endorsed or implemented in a major health system.

**Level 1B:** Annotation for a variant where the preponderance of evidence shows an association. The association must be replicated in more than one cohort with significant p-values, and preferably will have a strong effect size.

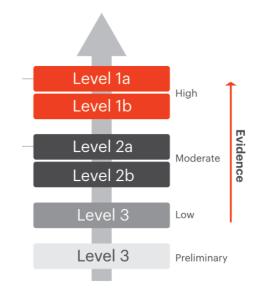
**Level 2A:** Annotation for a variant that qualifies for level 2B where the variant is within a Very Important known gene, so functional significance is more likely.

**Level 2B:** Annotation for a variant with moderate evidence of an association. The association must be replicated but there may be some studies that do not show statistical significance, and/or the effect size may be small.

**Level 3:** Annotation for a variant based on a single significant (not yet replicated) study or annotation for a variant evaluated in multiple studies but lacking clear evidence of an association.

**Level 4:** Annotation based on a case report, non-significant study or in vitro, molecular or functional assay evidence only.

Only genetic variants from level 1A to 2A were selected.



#### How was it analyzed?

The DNA was extracted from the buccal swab sample you provided and was analyzed by our clinical analysis laboratory. DNA was extracted using the KingFisher Flex® robotic extraction system (Thermo Fisher Scientific). The study of the genetic variants was performed by NGS (Next Generation Sequencing) using the Ion GeneStudio S5 system (Thermo Fisher Scientific).

### References





#### References

Scan the QR code to access our NutriGen report reference page.

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Together
we create the future
of personalizing medicine.





















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